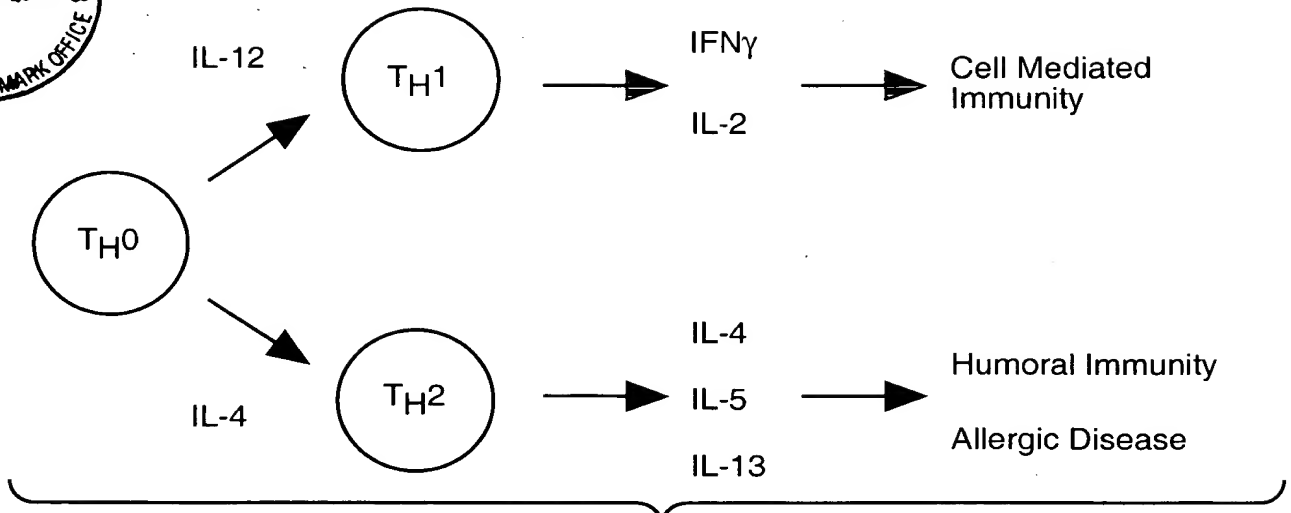
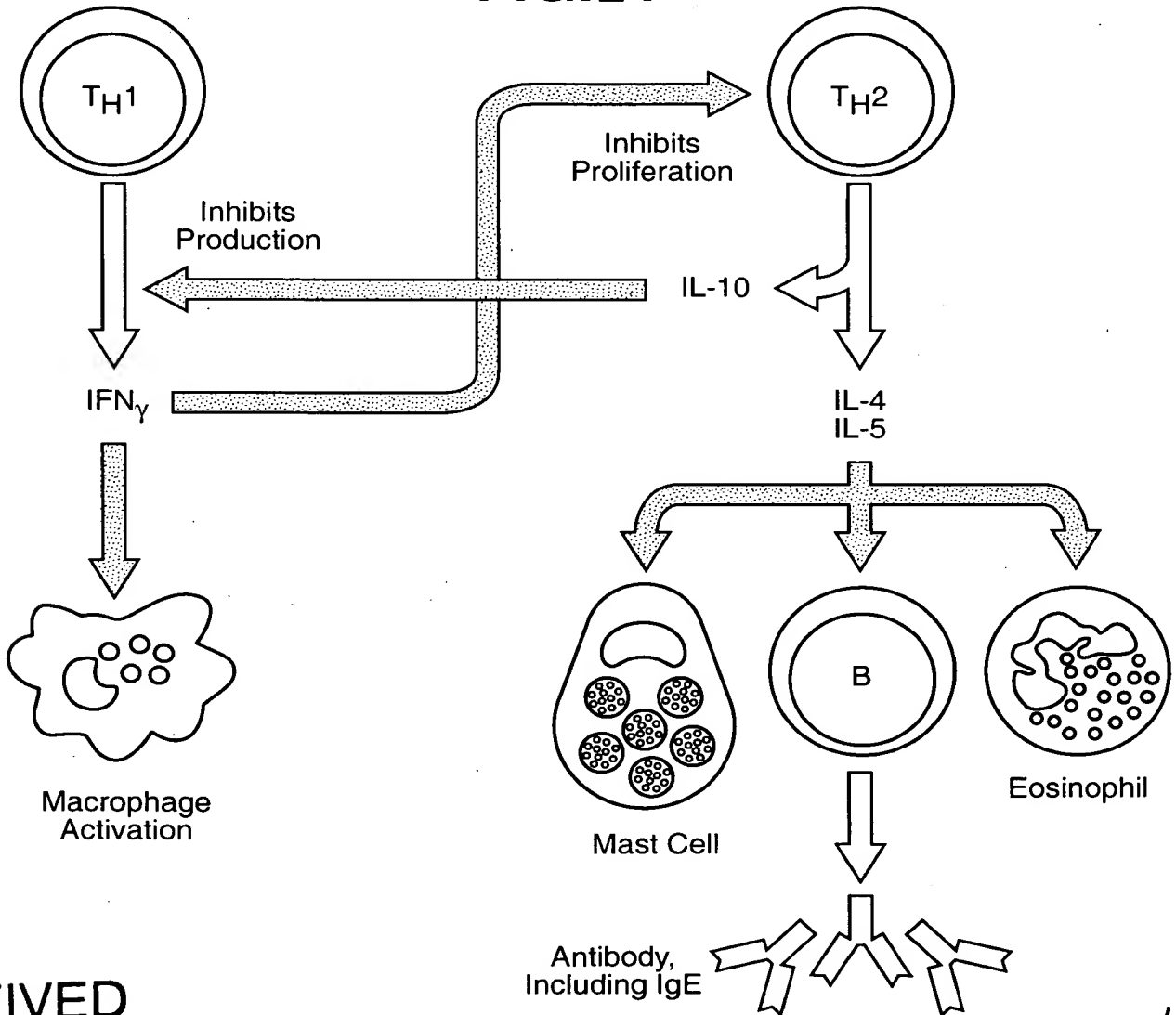




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**FIG. 1**



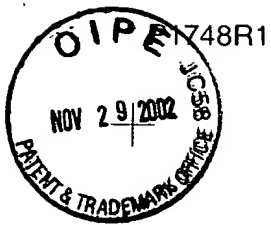
**FIG. 2**

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MRGGRGAPFWLWPLPKLALLPLLWVLFQRTRPQGSAGPLQCYGVGPLGDLNCSWEPLGD  
LGAPSELHLQSQKYRSNKTQTVAVAAGRSWVAIPREQLTMSDKLLVWGKAGQPLWPPV  
FVNLETQMKPNAPRLGPDVDFSEDDPLEATVHWAPPTWPSHKVLICQFHYRRCQEAAT  
LLEPELKTIPLTPEIQDLELATGYKVYGRCRMEKEEDLWGEWSPILSFQTPPSAPKDV  
WVSGNLCGTPGGEEPLLLWKAPGPCVQVSYKVWFVWGGRELSPEGITCCCSLIPSGAEW  
ARVSAVNATSWEPLTNLSLVCLDSASAPRSVAVSSIAGSTELLVTWQPGPGEPLEHVVD  
WARDGDPLEKLNWVRLPPGNLSALLPGNFTVGVPYRITVTAVSASGLASASSVWGFREE  
LAPLVGPTLWRLQDAPPGTPAIWAGEVPRHQLRGHLTHYTLCAQSGTSPSVCNMVSGNT  
QSVTLPLDLPWGPCELWVTASTIAGQGGPILRLHLPDNTLRWKVLPGLFLWGLFLLG  
CGLSLATSGRCYHLRHKVLPRVWWEKVPDPANSSSGQPHMEQVPEAQPLGDLPILEVEE  
MEPPPVMESSQPAQATAPLDSGYEKHFLPTPEELGLLGPPRPQVLA

APPROVED BY DRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS

**FIG.\_3**

MNRLRVARLTPLELLLSLMSLLLGTTRPHGSPGPLQCYSVGPLGILNCSWEPLGDLETTPV  
LYHQSQKYHPNRVWEVKVPSKQSWVTIPREQFTMADKLLIWGTQKGRPLWSSSVNLETQ  
MKPDTPQIFSQVDISEEATLEATVQWAPPVWPPQKALTCQFRYKECQAEAWTRLEPQLKT  
DGLTPVEMQNLEPGTCYQVSGRCQVENGYPWGEWSSPLSFQTPFLDPEDVWVSGTVCETS  
GKRAALLVWKDPRPCVQVTTYTVWFGAGDITTTQEEVPCCKSPVPAMMEWAVVSPGNSTSW  
VPPTNLSLVCLAPESAPCDVGVSSADGSPGIKVTWKQGTRKPLEYVVDWAQDGDSDKLN  
WTRLPPGNLSTLLPGEFKGGVPYRITVTAVYSGGLAAAPSVWGFREELVPLAGPAVWRLP  
DDPPGTPVVAWGEVPRHQLRGQATHYTFICIQSRGLSTVCRNVSSQTQTATLPNLHSGSFK  
LWVTVSTVAGQGPPGPDLSLHLPDNRIWKALPWFLSLWGLLLMGCGLSLASTRCLQARC  
LHWRHKLKPQWIWERPDPANSNSGQPYIKEVSLPQPPKDGPILEVEEVELQPVVESPKA  
SAPIYSGYEKHFLPTPEELGLLV

**FIG.\_4**

ADULT															FETAL					
PBLs	Colon	Sm. Int.	Ovary	Testis	Prostate	Thymus	Spleen	Heart	Brain	Placenta	Lung	Liver	Sk. Muscle	Kidney	Pancreas	Kidney	Liver	Lung	Brain	Heart

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**FIG.\_6**

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

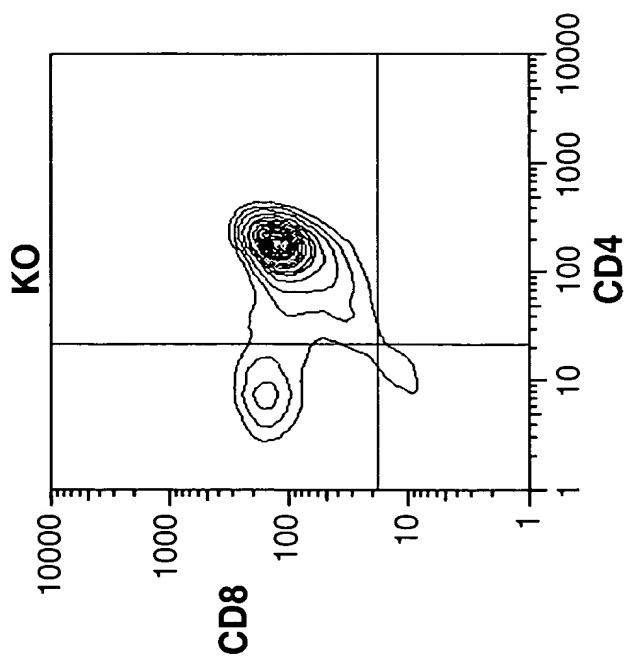
h-TCCR	1	MRGGRGAPFWLWLP	LPKLAALLPLLLWVLFQRTRP	QGSAGPLQCYGVGPLGDL
m-TCCR	1	-----MNRRLRVARI	TPLELLLSLSLSLLGTRPHGSP	GPLQCYSVGPLGLIL
h-TCCR	51	NCSWEPLGDLGAPSE	HLHLSQSKYRSNKTQTVA	AAAGRSWVAIPREQLTMS
m-TCCR	46	NCSWEPLGDLLET	PPVLYHQSKYHPNRVWEV	KPSKQSWVTIPREQFTMA
h-TCCR	101	DKLLVWGTAKAGQ	PLWPPVFNLEETQMKPNA	PRGLPDDVFSEDDPLEATVH
m-TCCR	96	DKLLIWGTQKGR	PLWSSVSVNLEETQMKPD	TPQIFSQVDISEEATLEATVQ
h-TCCR	151	WAPPTWPSHKVL	ICQFHYRRCCQEAATLLE	PELKTITPLTPVEIQDLELAT
m-TCCR	146	WAPPVWPPQKAL	ITCQFRYKECCQAEATRL	LEPQLKTDGLTPVEMQNLEPGT
h-TCCR	201	GKVVYGRCRMKEED	LWGEWSPILSFQTPPSA	PKDVWVSGNLCTPTGGEE
m-TCCR	195	CYQVSGRCQVENGY	P-WGEWSSPLSFQTPFLD	PEDEVVSGTVCETSGKRA
h-TCCR	251	PIILLWKA	PGPCVQVSQVYKVVFW	VGGRELSPEGITCCCSLIP
m-TCCR	245	ALLVWKDPRPCVQ	VTYTVWFAGDITTTQEE	VPEVPCCKSPVPAWMEWAVVSP
h-TCCR	301	VNATSWEP	LTNLSLVCLDSASAPRS	VAVSSIAGSTELLVTWQ
m-TCCR	295	GNSTSWVP	PTNLSLVCLAPESAPCD	VGVSSADGSPGIKVTWKQ
h-TCCR	351	HVVDWARDGD	PLEKLNWVRLPPGNLS	ALLPGNFTVGVFYRITVTAV
m-TCCR	345	YVVDWAQDGD	SLDKLNWTRLPPGNLS	TLLPGEFKGGVYRITVTAV
h-TCCR	401	LAASA	SSVWGFFREELAPLV	GPTLWRLQDAPPGT
m-TCCR	395	LAASA	PSVWGFFREELVPLA	GPVWRLPDDPPGT
h-TCCR	451	HYTLCAQ	SGTSPSVCMNVSGNT	QSVTLPLPWGPCE
m-TCCR	445	HYTFCIQ	SRGLSTVCRNVSSQ	TQTATLPNLSHSGS
h-TCCR	501	GPILLRLHLP	DNTRLRWKVLPGIL	FLWGLFLLCGGLSLATS
m-TCCR	495	GPDL	SLHLPDNRIRWKA	LPWFLSLWGLLLMCGGLSL
h-TCCR	547	HKVLP	PRWVWEKVPDPAN	SSSGQPHMEQVPEA
m-TCCR	545	HKLLPQ	WIWERVDPAN	SNVSGQPYIKEVSL
h-TCCR	597	MESSQ	PAQATAPLD	SGYEKHF
m-TCCR	595	VES	-----PKASAPI	YSGYEKHF

Box 1

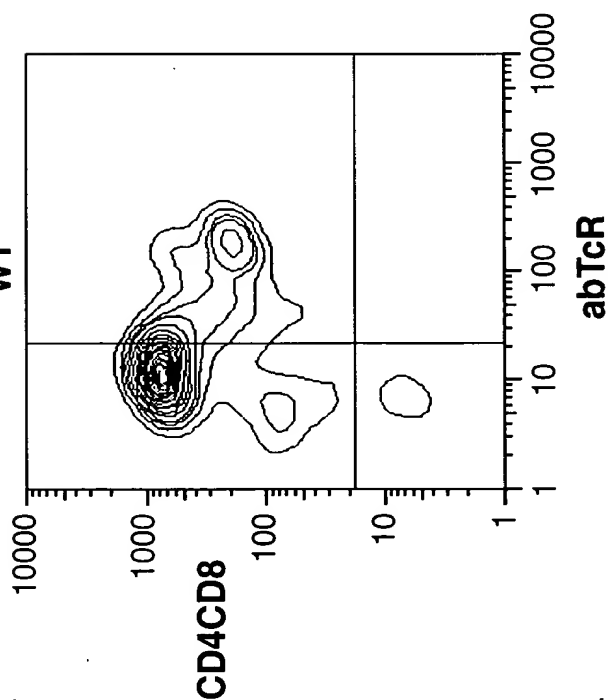
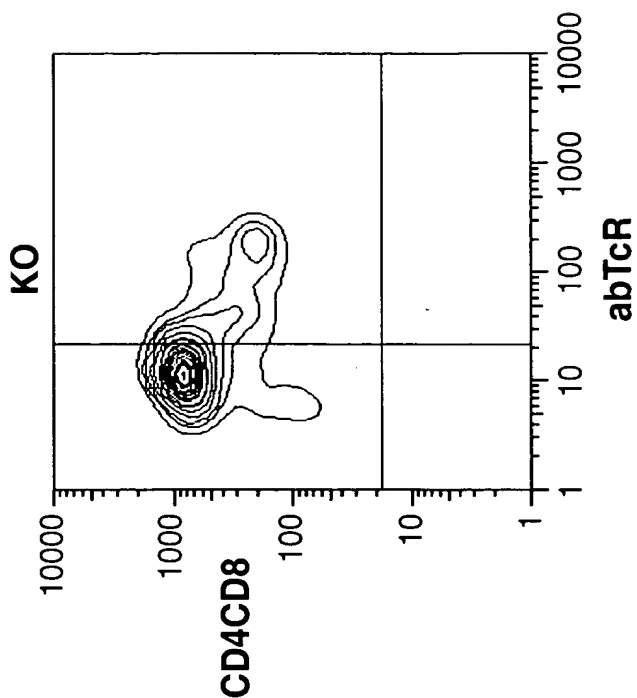
FIG.-5



APPROVED	O.G. FIG.
BY	CLASS
DRAFTSMAN	SUBCLASS

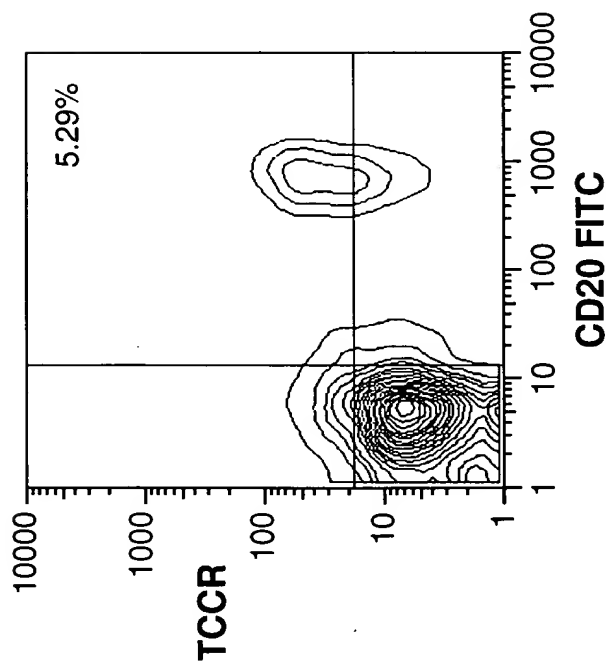
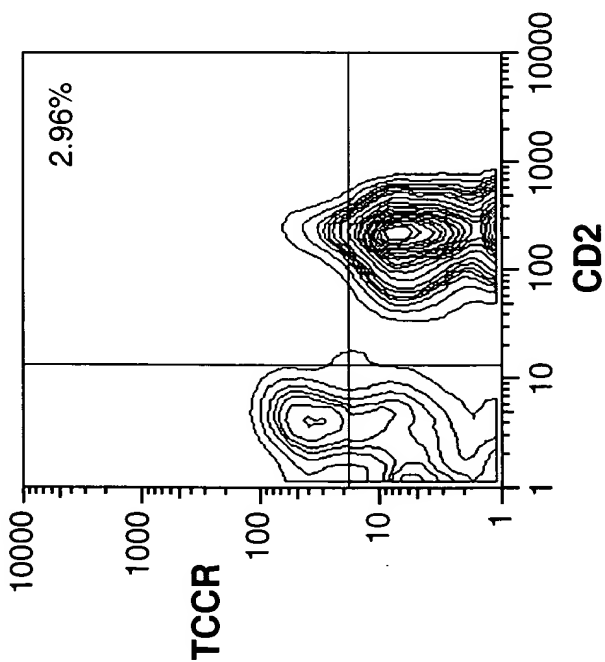
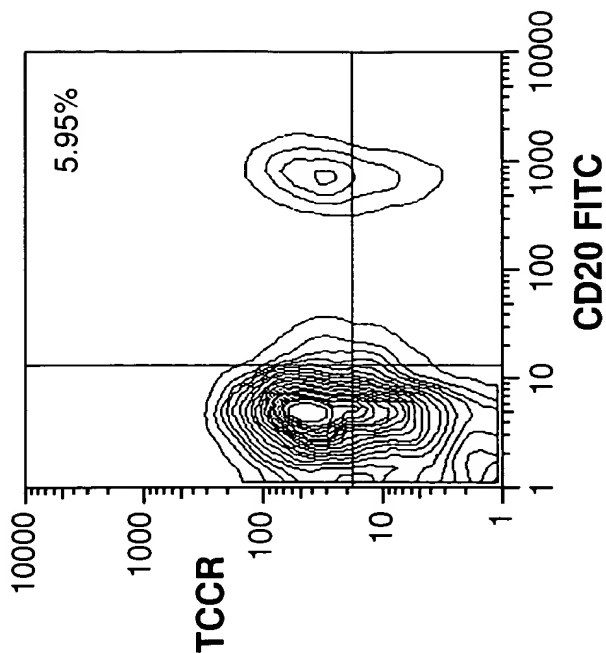
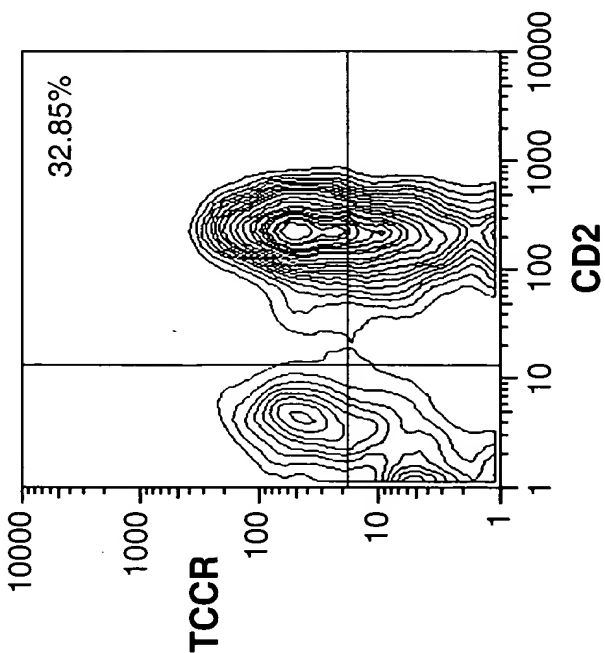


**FIG..7A**



**FIG..7B**

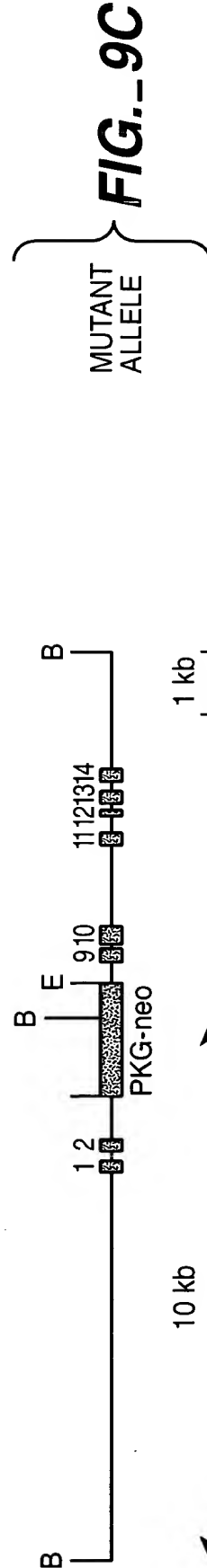
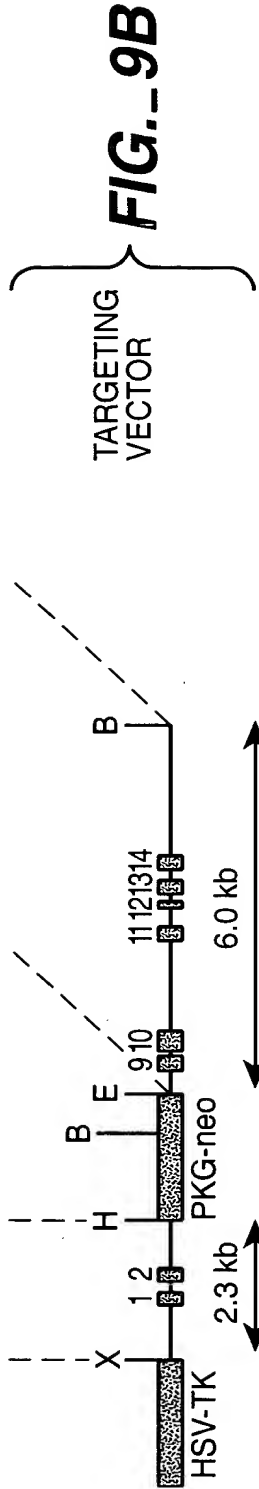
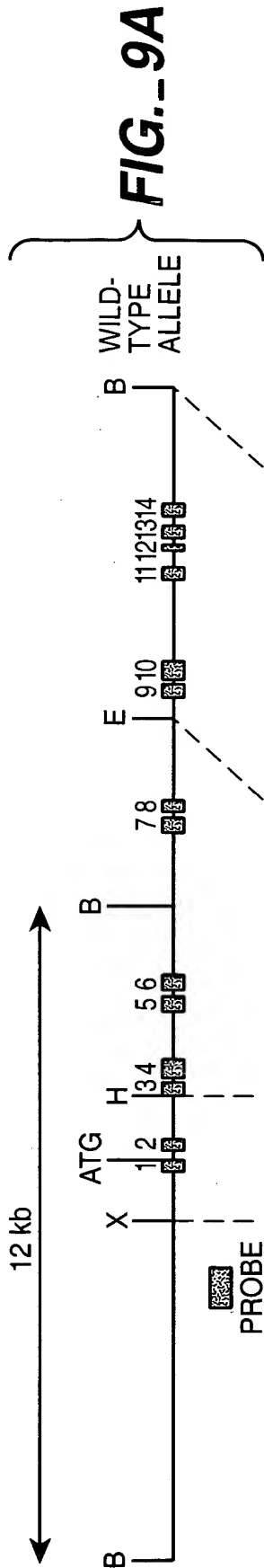
APPROVED BY DRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS



**FIG.-8A**

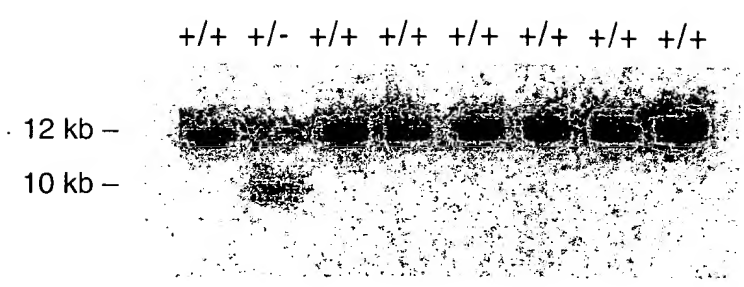
**FIG.-8B**

APPROVED BY DRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS

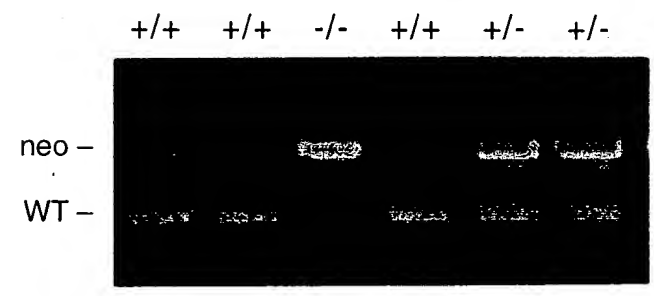


CLASS	SUBCLASS
BY	DRAFTSMAN

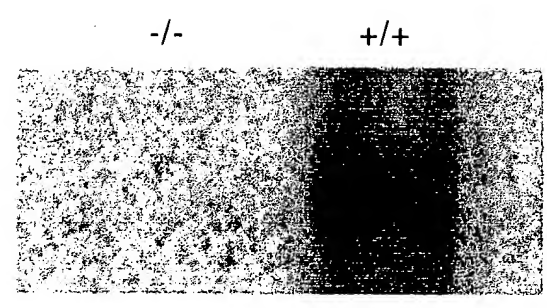
**FIG.\_10A**

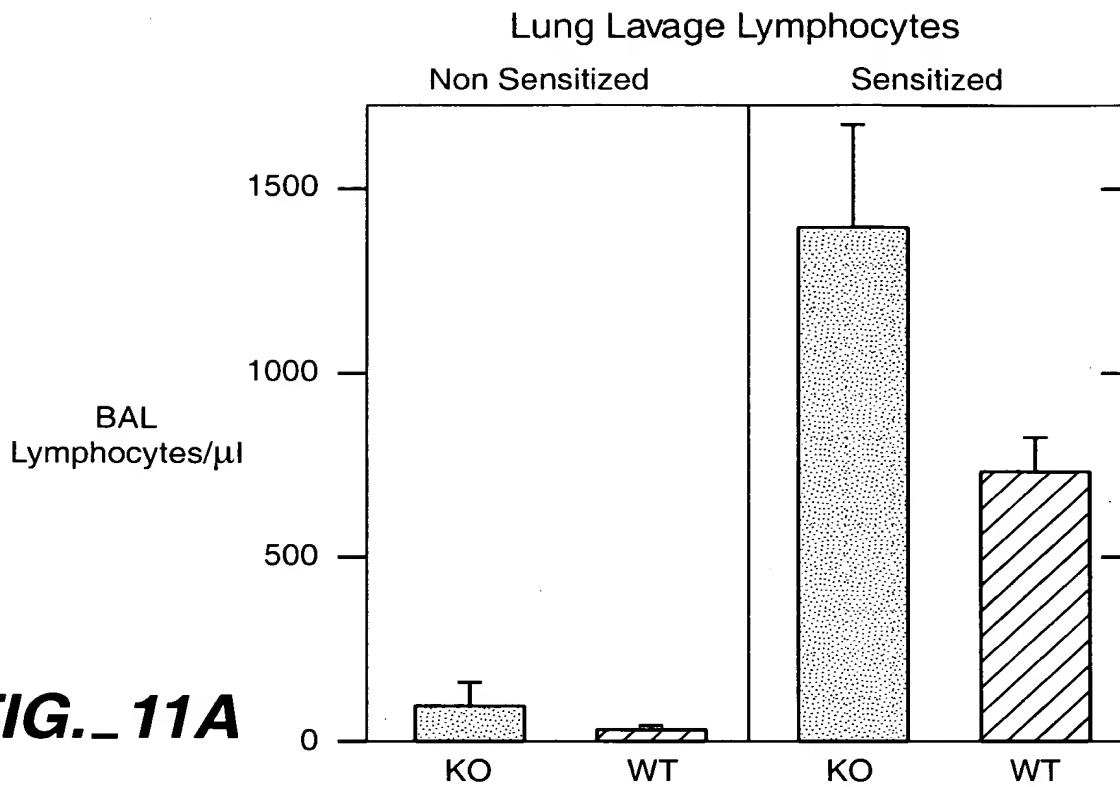


**FIG.\_10B**

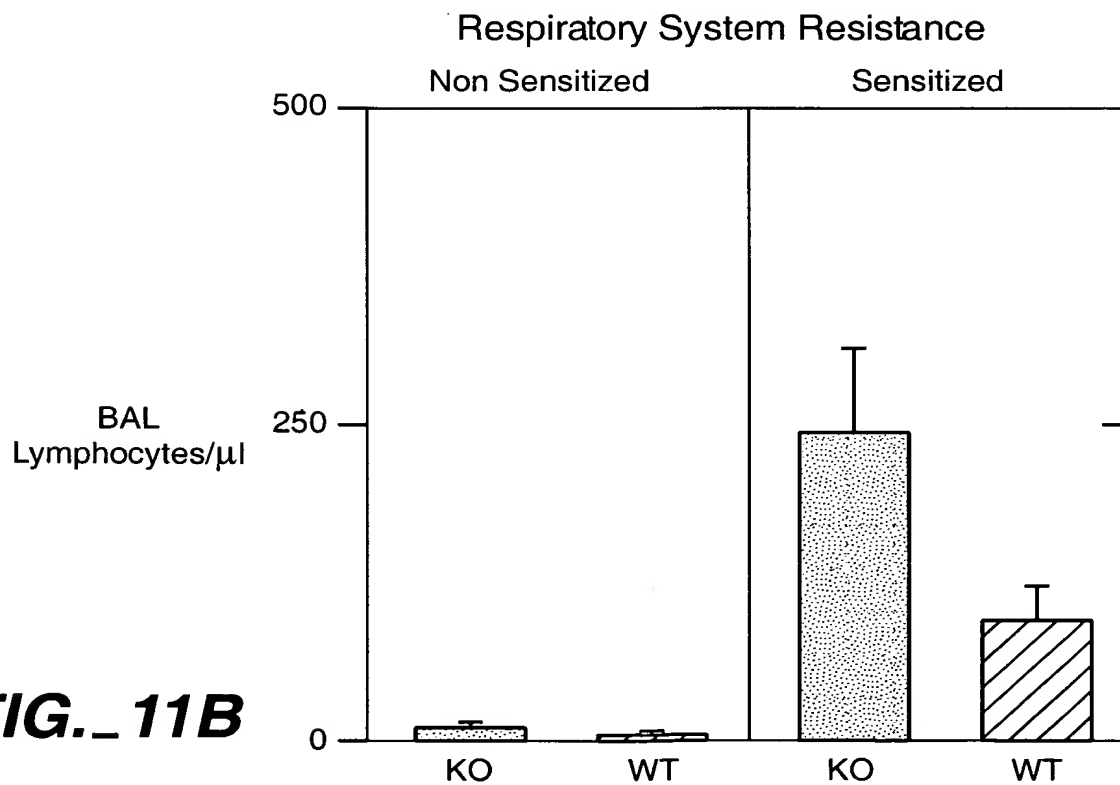


**FIG.\_10C**





**FIG.\_11A**

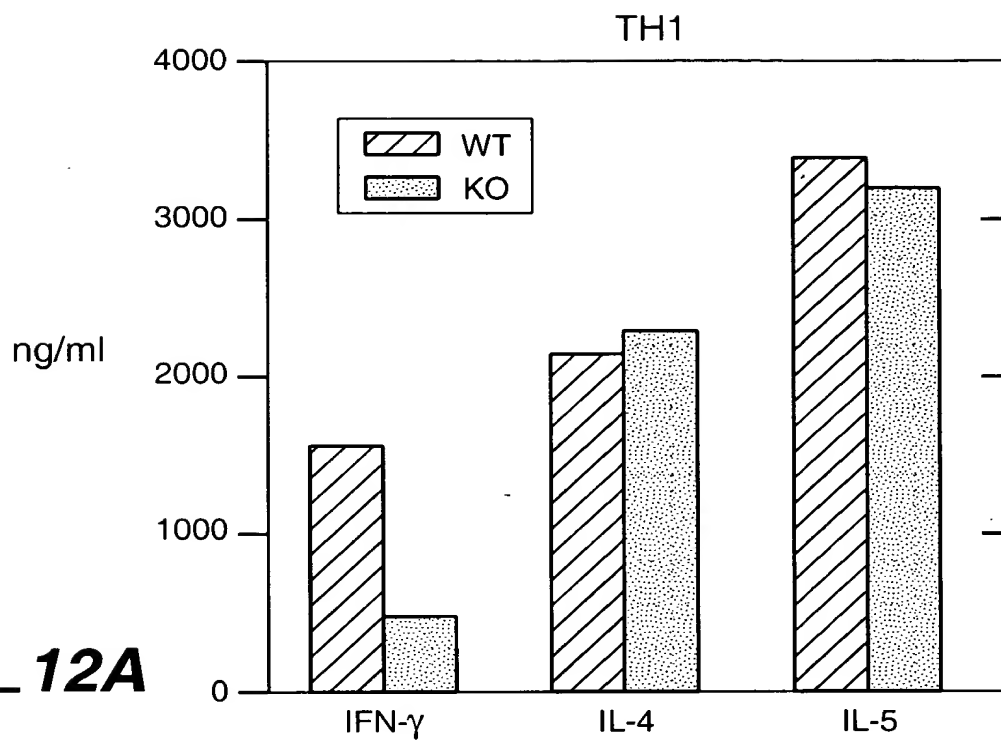


**FIG.\_11B**

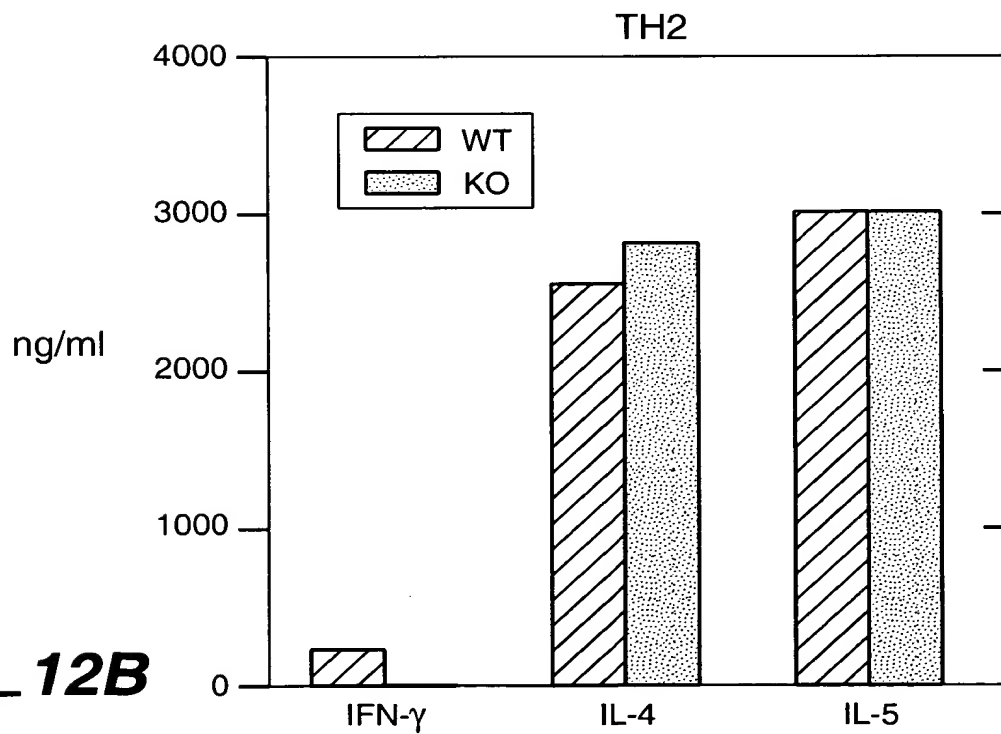


APPROVED	O.G. FIG.	SUBCLASS
BY	CLASS	
DRAFTSMAN		

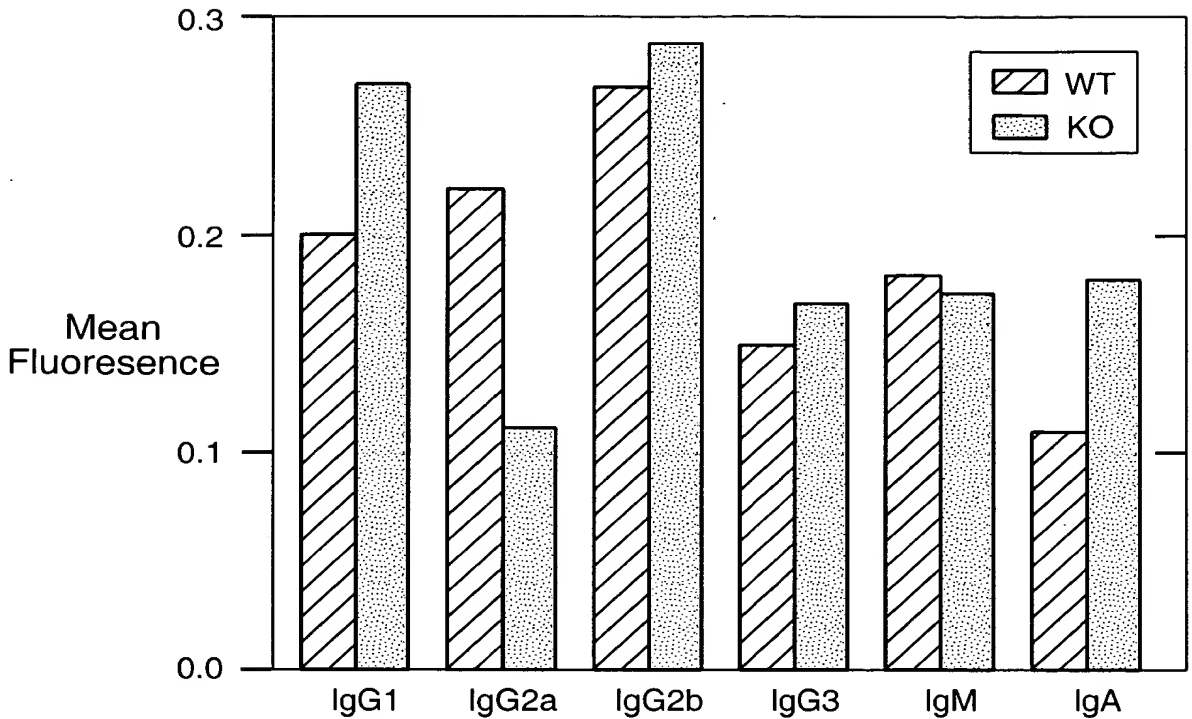
**FIG.\_12A**



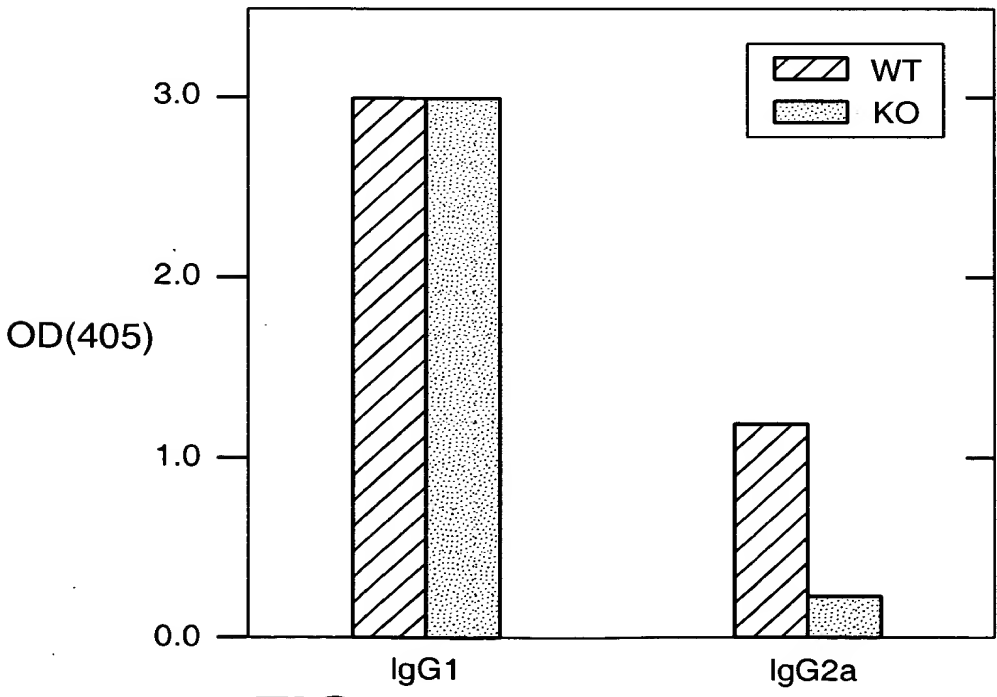
**FIG.\_12B**



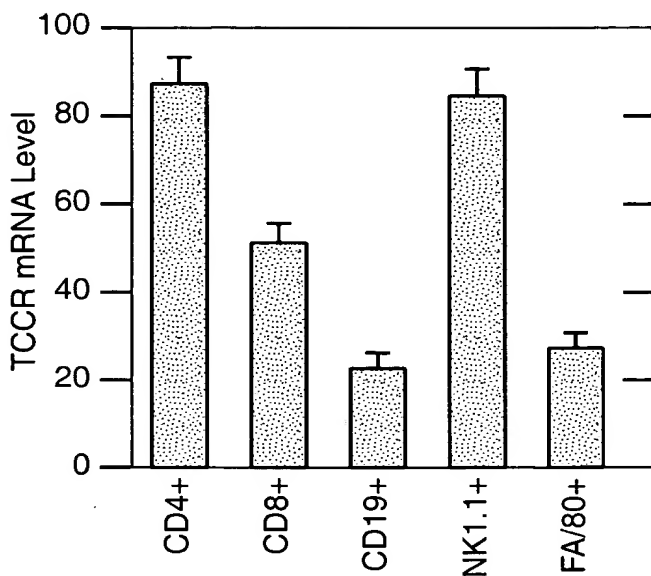
O.G. FIG.		SUBCLASS	
CLASS			
APPROVED BY DRAFTSMAN			



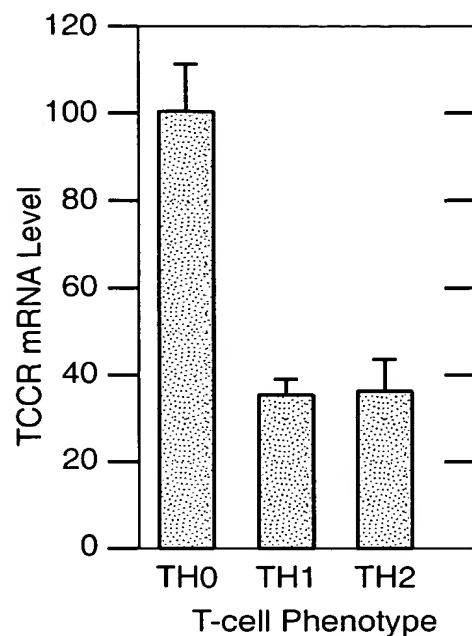
**FIG.\_13**



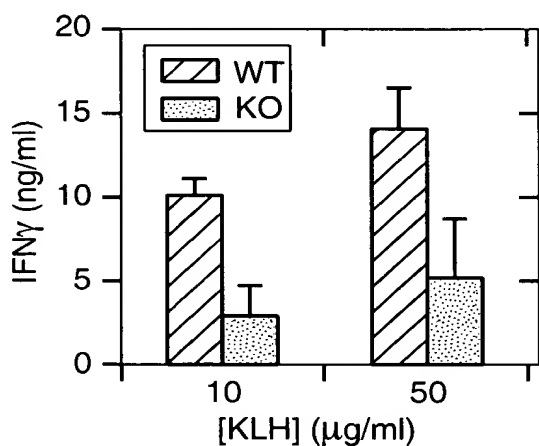
**FIG.\_14**



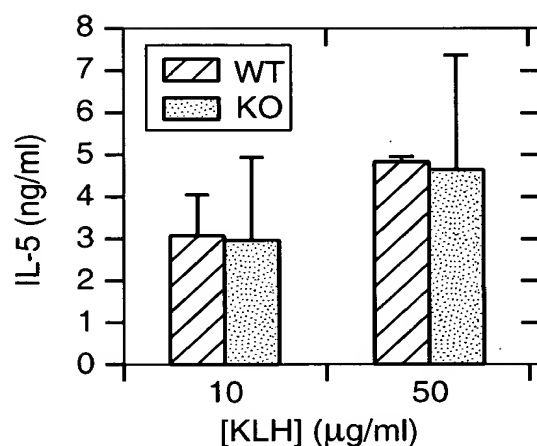
**FIG.\_15A**



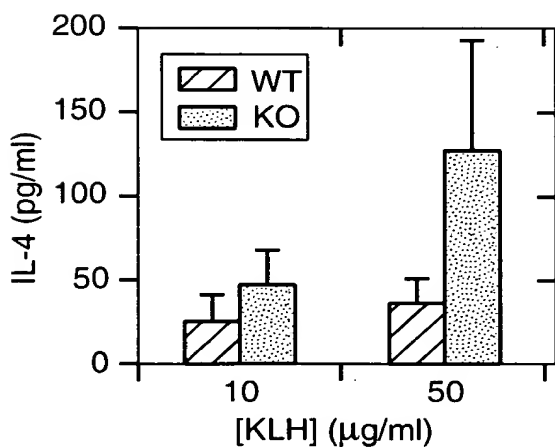
**FIG.\_15B**



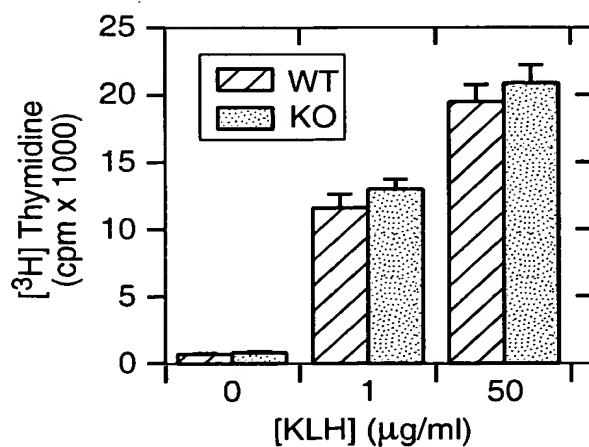
**FIG.\_16A**



**FIG.\_16C**



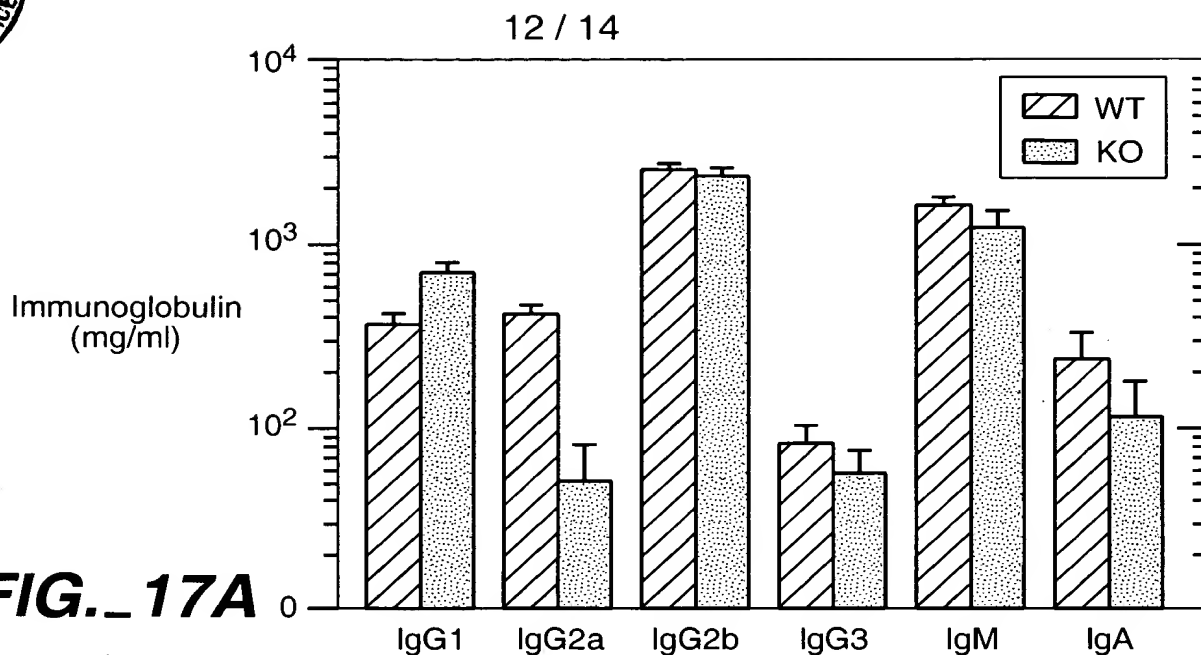
**FIG.\_16B**



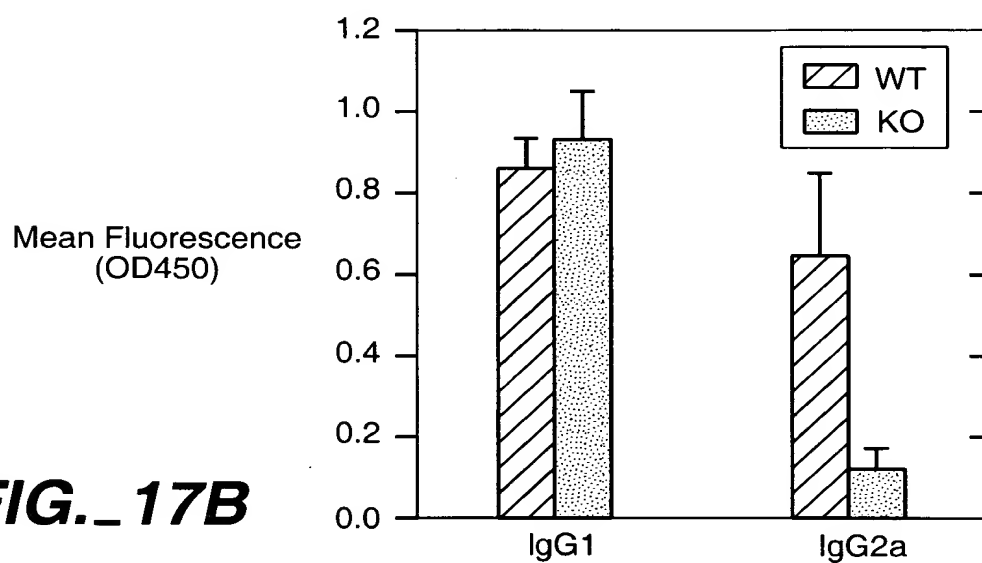
**FIG.\_16D**

APPROVED BY DRAFTSMAN	O.G. FIG.	SUBCLASS
	CLASS	

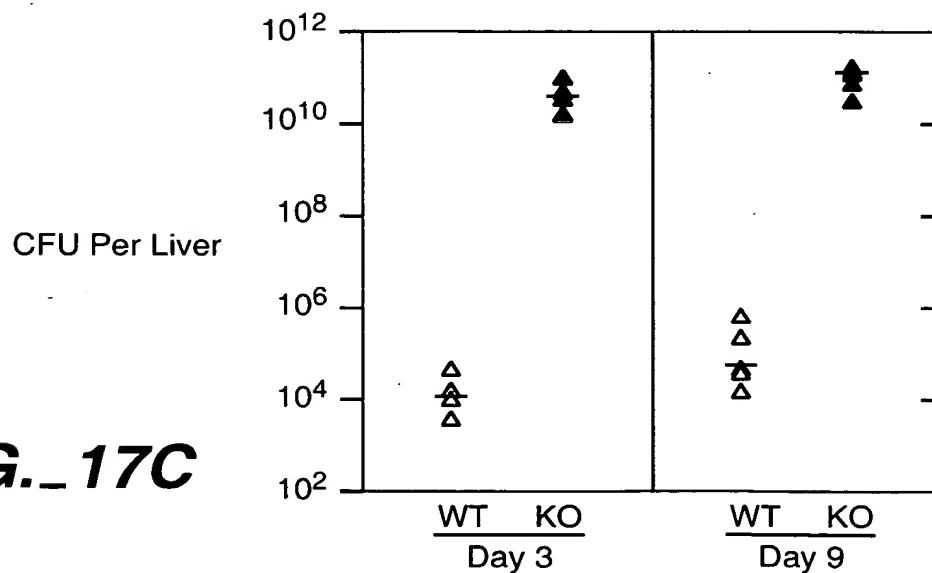
**FIG.\_17A**



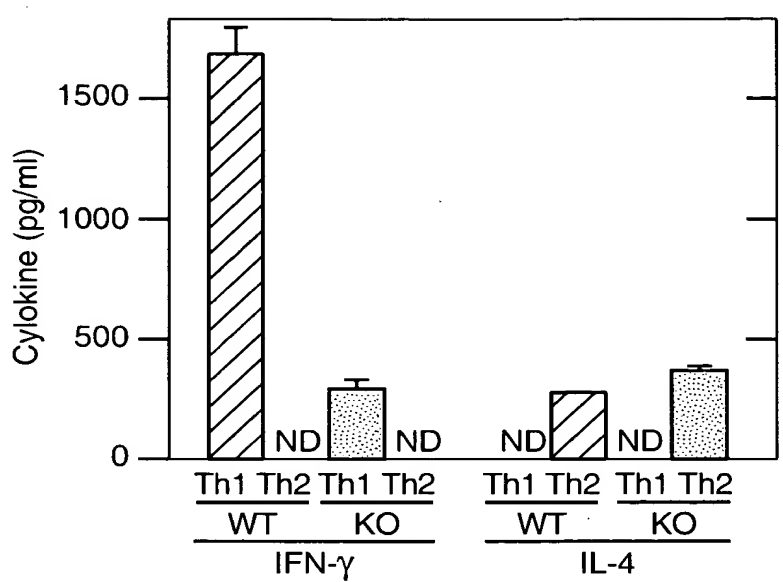
**FIG.\_17B**



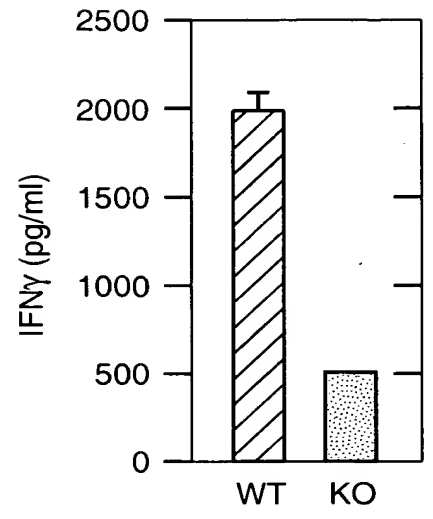
**FIG.\_17C**



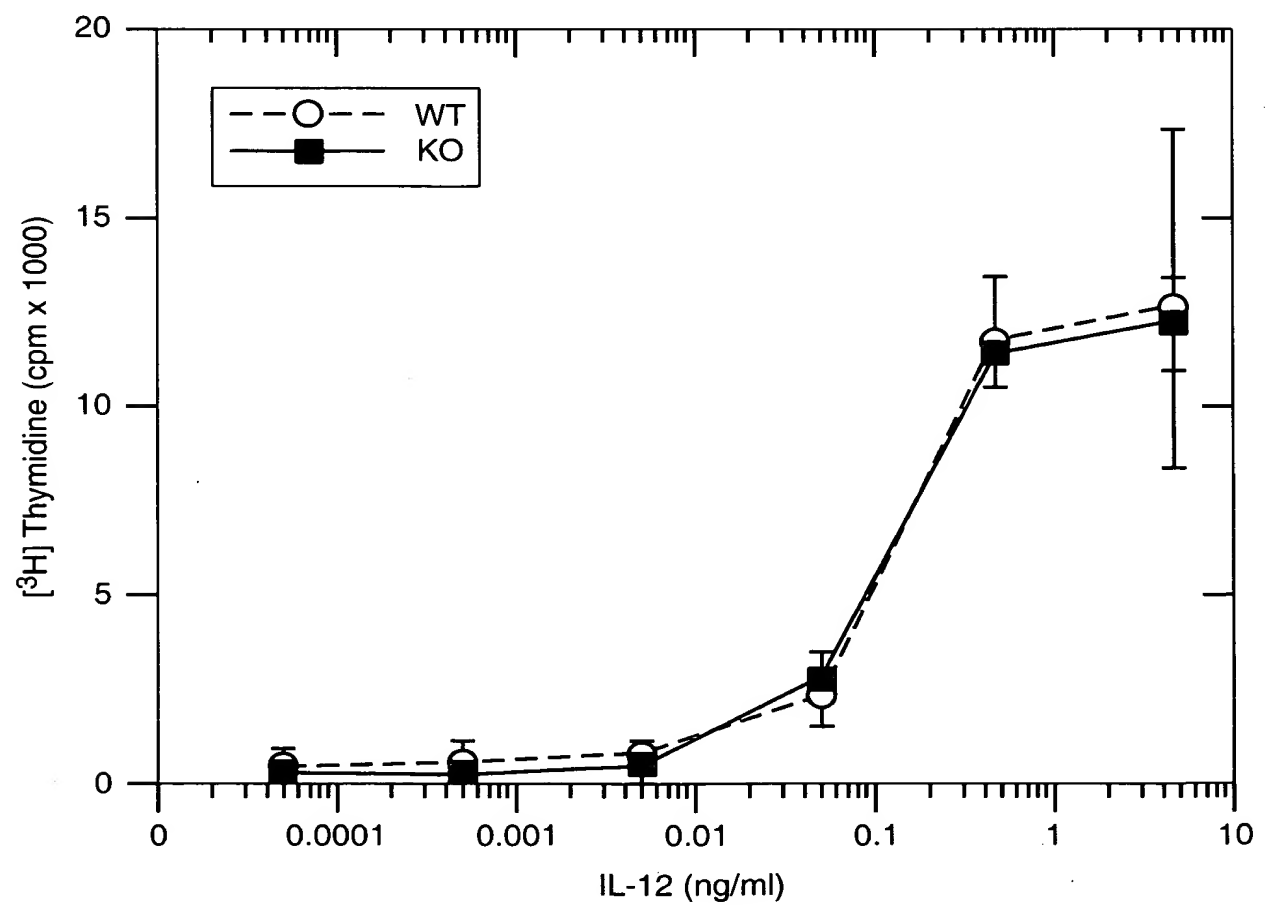
APPROVED BY DRAFTSMAN	O.G. FIG.	SUBCLASS
	CLASS	



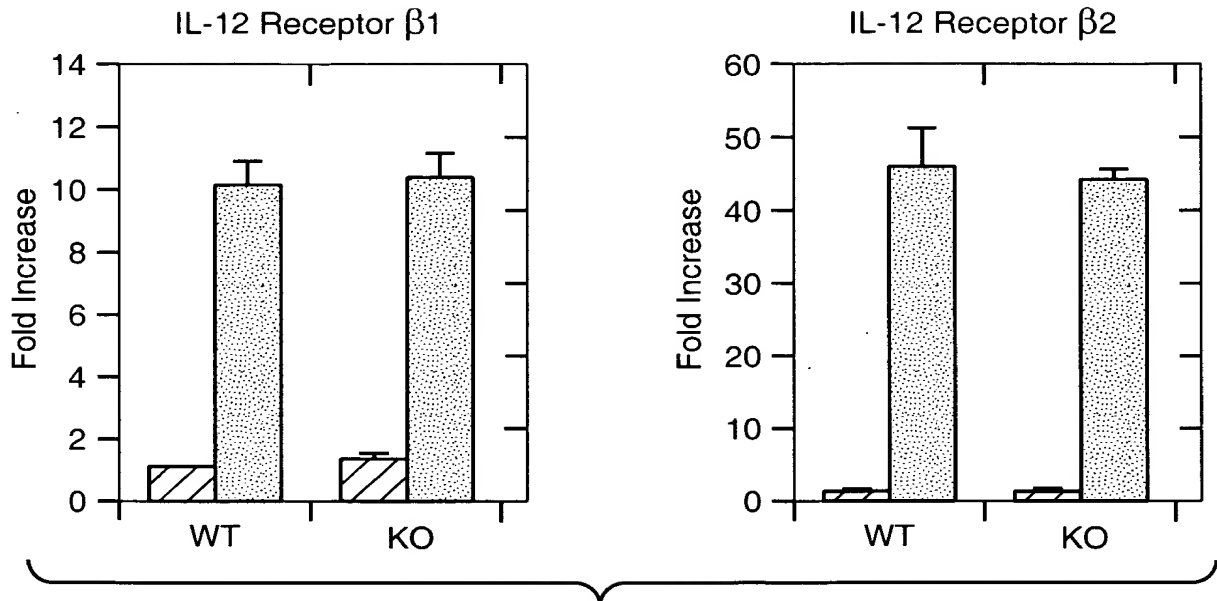
**FIG.\_18A**



**FIG.\_18B**



**FIG.\_18C**



**FIG.\_18D**

Primer / Probe	Sequence	SEQ ID NO:
mTCCR, sense, Taqman	TGGTCTCTCCTGGCAACAGC	5
mTCCR, as, Taqman	AGCCAAGCACACCAGAGACA	6
mTCCR, Taqman probe	CAGCTGGGTGCCTCCCACCAA	7
mRPL19, sense, Taqman	ATCCGCAAGCCTGTGACTGT	8
mRPL19, as, Taqman	TCGGGCCAGGGTGTTTTT	9
mRPL19, Taqman probe	TTCCCGGGCTCGTTGCCG	10
mIL12Rb1, sense, Taqman	TCGCGTCTCTGGGAAGCT	11
mIL12Rb1, as, Taqman	TTTAAGCCAATGTATCCGAGACTG	12
mIL12Rb1, Taqman probe	CGCCAGCGTCCTCCTCGTGG	13
mIL12Rb2, sense, Taqman	CAAGCATTTGCATCGCTATCA	14
mIL12Rb2, as, Taqman	AATGCCTTTTGCCGGAAGT	15
mIL12Rb2, Taqman probe	ACGAATTGAGAACGTGCCCACCGT	16

**FIG.\_19**